

RAW SEQUENCE LISTING

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Application Serial Number: 10/780,507
Source: IFWO
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DATE: 11/26/2004

PATENT APPLICATION: US/10/780,507

TIME: 10:21:34

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Output Set: N:\CRF4\11262004\J780507.raw

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3 <110> APPLICANT: MULLINS, James I.
4   RODRIGO, Allen G.
5   LEARN, Gerald H.
6   LI, Fusheng
7   NICKLE, David C.
8   JENSEN, Mark A.
10 <120> TITLE OF INVENTION: ANCESTRAL AND COT VIRAL SEQUENCES, PROTEINS AND IMMUNOGENIC
COMPOSITIONS
12 <130> FILE REFERENCE: 16336-001320US
14 <140> CURRENT APPLICATION NUMBER: 10/780,507
15 <141> CURRENT FILING DATE: 2004-02-17
17 <150> PRIOR APPLICATION NUMBER: US 10/204,204
18 <151> PRIOR FILING DATE: 2001-02-16
20 <150> PRIOR APPLICATION NUMBER: PCT/US01/05288
21 <151> PRIOR FILING DATE: 2001-02-16
23 <150> PRIOR APPLICATION NUMBER: US 60/183,659
24 <151> PRIOR FILING DATE: 2000-02-18
26 <150> PRIOR APPLICATION NUMBER: US 60/447,586
27 <151> PRIOR FILING DATE: 2003-02-14
29 <160> NUMBER OF SEQ ID NOS: 125
31 <170> SOFTWARE: PatentIn version 3.1
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34 <211> LENGTH: 2652
35 <212> TYPE: DNA
36 <213> ORGANISM: Artificial Sequence
38 <220> FEATURE:
39 <223> OTHER INFORMATION: Ancestral HIV-1 group M, subtype B, env sequence
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134 <211> LENGTH: 883

135 <212> TYPE: PRT

136 <213> ORGANISM: Artificial Sequence

138 <220> FEATURE:

139 <223> OTHER INFORMATION: Ancestral HIV-1 group M, subtype B, env sequence

141 <400> SEQUENCE: 2

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148           20           25           30
151 Leu Trp Val Thr Val Tyr Tyr Gly Val Pro Val Trp Lys Glu Ala Thr
152           35           40           45
155 Thr Thr Leu Phe Cys Ala Ser Asp Ala Lys Ala Tyr Asp Thr Glu Val
156           50           55           60
159 His Asn Val Trp Ala Thr His Ala Cys Val Pro Thr Asp Pro Asn Pro
160 65           70           75           80
163 Gln Glu Val Val Leu Glu Asn Val Thr Glu Asn Phe Asn Met Trp Lys
164           85           90           95
167 Asn Asn Met Val Glu Gln Met His Glu Asp Ile Ile Ser Leu Trp Asp

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175	Asn	Cys	Thr	Asp	Asp	Leu	Arg	Thr	Asn	Ala	Thr	Asn	Thr	Thr	Asn	Ser
176		130					135						140			
179	Ser	Ala	Thr	Thr	Asn	Thr	Thr	Ser	Ser	Gly	Gly	Gly	Thr	Met	Glu	Gly
180	145					150					155					160
183	Glu	Lys	Gly	Glu	Ile	Lys	Asn	Cys	Ser	Phe	Asn	Val	Thr	Thr	Ser	Ile
184					165					170					175	
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212			275					280					285			
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220	305					310					315					320
223	Arg	Lys	Ser	Ile	Pro	Ile	Gly	Pro	Gly	Arg	Ala	Leu	Tyr	Ala	Thr	Gly
224					325					330					335	
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231	Lys	Trp	Asn	Asn	Thr	Leu	Lys	Gln	Ile	Val	Thr	Lys	Leu	Arg	Glu	Gln
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240	385					390					395					400
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271 Lys Val Val Lys Ile Glu Pro Leu Gly Val Ala Pro Thr Lys Ala Lys
272           515           520           525
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276           530           535           540
279 Met Phe Leu Gly Phe Leu Gly Ala Ala Gly Ser Thr Met Gly Ala Ala
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283 Ser Met Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val
284           565           570           575
287 Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu
288           580           585           590
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292           595           600           605
295 Ala Val Glu Arg Tyr Leu Lys Asp Gln Gln Leu Leu Gly Ile Trp Gly
296           610           615           620
299 Cys Ser Gly Lys Leu Ile Cys Thr Thr Ala Val Pro Trp Asn Ala Ser
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303 Trp Ser Asn Lys Ser Leu Asp Lys Ile Trp Asn Asn Met Thr Trp Met
304           645           650           655
307 Glu Trp Glu Arg Glu Ile Asp Asn Tyr Thr Gly Leu Ile Tyr Thr Leu
308           660           665           670
311 Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu Leu Leu
312           675           680           685
315 Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe Asp Ile Thr Asn
316           690           695           700
319 Trp Leu Trp Tyr Ile Lys Ile Phe Ile Met Ile Val Gly Gly Leu Val
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327 Gln Gly Tyr Ser Pro Leu Ser Phe Gln Thr Arg Leu Pro Ala Pro Arg
328           740           745           750
331 Gly Pro Asp Arg Pro Glu Gly Ile Glu Glu Glu Gly Gly Glu Arg Asp
332           755           760           765
335 Arg Asp Arg Ser Gly Arg Leu Val Asn Gly Phe Leu Ala Leu Ile Trp
336           770           775           780
339 Asp Asp Leu Arg Ser Leu Cys Leu Phe Ser Tyr His Arg Leu Arg Asp
340 785           790           795           800
343 Leu Leu Leu Ile Val Ala Arg Ile Val Glu Leu Leu Gly Arg Arg Gly
344           805           810           815
347 Trp Glu Ala Leu Lys Tyr Trp Trp Asn Leu Leu Gln Tyr Trp Ser Gln
348           820           825           830
351 Glu Leu Lys Asn Ser Ala Val Ser Leu Leu Asn Ala Thr Ala Ile Ala
352           835           840           845
355 Val Ala Glu Gly Thr Asp Arg Val Ile Glu Val Val Gln Arg Ala Cys
356           850           855           860
359 Arg Ala Ile Leu His Ile Pro Arg Arg Ile Arg Gln Gly Leu Glu Arg
360 865           870           875           880
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369 <212> TYPE: DNA
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372 <220> FEATURE:
373 <223> OTHER INFORMATION: Ancestral HIV-1 group M, subtype C, env sequence
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VERIFICATION SUMMARY

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